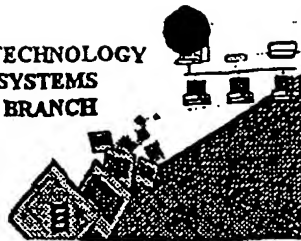




RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10/775,487
FFMO
7-16-04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENT/IN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/efc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Revised 05/17/04

Raw Sequence Listing Error Summary

Best Available Copy

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/775,487

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
Numbering The numbering under each 3rd amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5 Variable Length Sequence(s) _____ contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>.<223> section that some may be missing.
- 6 PatentIn 2.0
"bug" A "bug" in PatentIn version 2.0 has caused the <220>.<223> section to be missing from amino acid sequence(s) _____. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>.<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>.<223> sections for Artificial or Unknown sequences.
- 7 Skipped Sequences
(OLD RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
(2) INFORMATION FOR SEQ ID NO X (insert SEQ ID NO where "X" is shown)
(i) SEQUENCE CHARACTERISTICS (Do not insert any subheadings under this heading)
(xi) SEQUENCE DESCRIPTION SEQ ID NO X (insert SEQ ID NO where "X" is shown)
This sequence is intentionally skipped.

Please also adjust the "(ii) NUMBER OF SEQUENCES" response to include the skipped sequences.
- 8 Skipped Sequences
(NEW RULES) Sequence(s) _____ missing. If intentional, please insert the following lines for each skipped sequence:
<210> sequence id number
<400> sequence id number
000
- 9 Use of n's or Xaa's
(NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
Per 1.823 of Sequence Rules, use of <220>.<223> is MANDATORY if n's or Xaa's are present.
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <21>
Response Per 1.823 of Sequence Rules, the only valid <21> responses are Unknown, Artificial Sequence, or Scientific name (Genus/species). <220>.<223> section is required when <21> response is Unknown or Artificial Sequence.
- 11 Use of <220> Sequence(s) _____ missing the <220> "Feature" and associated numeric identifiers and responses.
Use of <220> to <223> is MANDATORY if <21> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
(See "Federal Register," 01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
"bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid.



IFWO

RAW SEQUENCE LISTING

DATE: 12/16/2004

PATENT APPLICATION: US/10/775,487

TIME: 15:24:13

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\12162004\J775487.raw

4 <110> APPLICANT: Faustman, Denise L.
 5 Hayashi, Takuma
 7 <120> TITLE OF INVENTION: Methods For Diagnosing Autoimmune
 8 Disease
 10 <130> FILE REFERENCE: 00786/457003
 12 <140> CURRENT APPLICATION NUMBER: 10/775,487
 13 <141> CURRENT FILING DATE: 2004-02-10
 15 <150> PRIOR APPLICATION NUMBER: 09/258,682
 16 <151> PRIOR FILING DATE: 1999-02-26
 18 <150> PRIOR APPLICATION NUMBER: 09/031,629
 19 <151> PRIOR FILING DATE: 1998-02-27
 21 <160> NUMBER OF SEQ ID NOS: 11
 23 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 25 <210> SEQ ID NO: 1
 26 <211> LENGTH: 32
 27 <212> TYPE: DNA
 28 <213> ORGANISM: Genus/Species
 30 <400> SEQUENCE: 1
 31 gatctagga cttccgctg gggactttcc ag
 33 <210> SEQ ID NO: 2
 34 <211> LENGTH: 40
 35 <212> TYPE: DNA
 36 <213> ORGANISM: genus/species
 38 <400> SEQUENCE: 2
 39 gatctcaggg gaatctcct ctccttttat gggcgtagcg
 41 <210> SEQ ID NO: 3
 42 <211> LENGTH: 10
 43 <212> TYPE: DNA
 44 <213> ORGANISM: genus/species
 46 <400> SEQUENCE: 3
 47 gggactttcc
 49 <210> SEQ ID NO: 4
 50 <211> LENGTH: 10
 51 <212> TYPE: DNA
 52 <213> ORGANISM: genus species
 54 <400> SEQUENCE: 4
 55 gggacgtccc
 57 <210> SEQ ID NO: 5
 58 <211> LENGTH: 10
 59 <212> TYPE: DNA
 60 <213> ORGANISM: genus species
 62 <400> SEQUENCE: 5
 63 ggaatttcc

Invalid
ResponseInvalid
ResponseSame
errorSame
errorSame
error

(pg. 1-2)

← Mandatory
 response
 (213)³² has to
 be Artificial
 Unknown or
 a Genus
 or Species

see item
 #10 on
 error Summary
 Sheet.

FYI: which genus/
 species?

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/775,487

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Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\12162004\J775487.raw

65 <210> SEQ ID NO: 6
 66 <211> LENGTH: 5
 67 <212> TYPE: DNA
 68 <213> ORGANISM: genus species
 70 <400> SEQUENCE: 5
 71 gggac
 73 <210> SEQ ID NO: 7
 74 <211> LENGTH: 5
 75 <212> TYPE: DNA
 76 <213> ORGANISM: genus species
 78 <400> SEQUENCE: 7
 79 ggaaa
 81 <210> SEQ ID NO: 8
 82 <211> LENGTH: 5
 83 <212> TYPE: DNA
 84 <213> ORGANISM: genus species
 86 <400> SEQUENCE: 8
 87 ctcac
 89 <210> SEQ ID NO: 9
 90 <211> LENGTH: 5
 91 <212> TYPE: DNA
 92 <213> ORGANISM: genus species
 94 <400> SEQUENCE: 9
 95 tttcc
 97 <210> SEQ ID NO: 10
 98 <211> LENGTH: 5
 99 <212> TYPE: DNA
 100 <213> ORGANISM: genus species
 102 <400> SEQUENCE: 10
 103 gtccc
 105 <210> SEQ ID NO: 11
 106 <211> LENGTH: 5
 107 <212> TYPE: DNA
 108 <213> ORGANISM: genus species
 110 <400> SEQUENCE: 11
 111 tttcc

Same
errors

Which genus/
species?

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/775,487

DATE: 12/16/2004

TIME: 15:24:14

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\12162004\J775487.raw

Best Available Copy